



SEQUENCE LISTING

<10> Nelsestuen, Gary L.

<120> MODIFIED VITAMIN K-DEPENDENT POLYPEPTIDES

<130> 09531-016002

<140> US 10/031,005

<141> 2001-10-29

<150> PCT/US00/11416

<151> 2000-04-28

<150> US 09/302,239

<151> 1999-04-29

<150> US 08/955,636

<151> 1997-10-23

<160> 21

<170> FastSEQ for Windows Version 4.0

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<212> PRT

<213> Homo sapiens

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<222> (1)...(44)

<223> Xaa = gamma carboxyglutamic acid or glutamic acid

<400> 1

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|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ala | Asn | Ser | Phe | Leu | Xaa | Xaa | Leu | Arg | His | Ser | Ser | Leu | Xaa | Arg | Xaa |
| 1 | | | | 5 | | | | 10 | | | | | 15 | | |
| Cys | Ile | Xaa | Xaa | Ile | Cys | Asp | Phe | Xaa | Xaa | Ala | Lys | Xaa | Ile | Phe | Gln |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| Asn | Val | Asp | Asp | Thr | Leu | Ala | Phe | Trp | Ser | Lys | His | | | | |
| | | 35 | | | | | 40 | | | | | | | | |

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<400> 2

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|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ala | Asn | Ser | Phe | Leu | Xaa | Xaa | Leu | Arg | Pro | Gly | Asn | Val | Xaa | Arg | Xaa |
| 1 | | | | 5 | | | | 10 | | | | | 15 | | |
| Cys | Ser | Xaa | Xaa | Val | Cys | Xaa | Phe | Xaa | Xaa | Ala | Arg | Xaa | Ile | Phe | Gln |

20 25 30
 Asn Thr Xaa Asp Thr Met Ala Phe Trp Ser Phe Tyr
 35 40

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<400> 3
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 1 5 10 15
 Cys Lys Xaa Xaa Gln Cys Ser Phe Xaa Xaa Ala Arg Xaa Ile Phe Lys
 20 25 30
 Asp Ala Xaa Arg Thr Lys Leu Phe Trp Ile Ser Tyr
 35 40

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<400> 4
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 1 5 10 15
 Cys Arg Xaa Xaa Leu Cys Ser Phe Xaa Xaa Ala His Xaa Ile Phe Arg
 20 25 30
 Asn Xaa Xaa Arg Thr Arg Gln Phe Trp Val Ser Tyr
 35 40

<210> 5
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<400> 5
 Tyr Asn Ser Gly Lys Leu Xaa Xaa Phe Val Gln Gly Asn Leu Xaa Arg
 1 5 10 15
 Xaa Cys Met Xaa Xaa Lys Cys Ser Phe Xaa Xaa Ala Arg Xaa Val Phe
 20 25 30
 Xaa Asn Thr Xaa Arg Thr Thr Xaa Phe Trp Lys Gln Tyr
 35 40 45

<210> 6
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 1 5 10 15
 Xaa Cys Met Xaa Xaa Lys Cys Ser Phe Xaa Xaa Ala Arg Xaa Val Phe
 20 25 30
 Xaa Asn Thr Xaa Lys Arg Thr Thr Xaa Phe Trp Lys Gln Tyr
 35 40 45

<210> 7
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 <212> DNA
 <213> Artificial Sequence

<220>
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 aaattaatac gactcactat agggagaccc aagctt 36

<210> 8
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<400> 8
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<400> 9
 acgctccacg ttgccgtgcc gcagctctc taggaa 36

<210> 10
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<400> 15
cagtgtgtca tccacatttt cgaaaatttc cttggc

36

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36

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Xaa Cys Leu Xaa Xaa Pro Cys Ser Arg Xaa Xaa Ala Phe Xaa Ala Leu
20 25 30
Xaa Ser Leu Ser Ala Thr Asp Ala Phe Trp Ala Lys Tyr
35 40 45

<210> 18
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<212> PRT
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<400> 18
Ala Asn Ser Phe Leu Xaa Xaa Val Lys Gln Gly Asn Leu Xaa Arg Xaa
1 5 10 15
Cys Leu Xaa Xaa Ala Cys Ser Leu Xaa Xaa Ala Arg Xaa Val Phe Xaa
20 25 30
Asp Ala Xaa Gln Thr Asp Xaa Phe Trp Ser Lys Tyr
35 40

<210> 19
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<221> MOD_RES
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<400> 19
Ala Asn Ser Leu Leu Xaa Xaa Thr Lys Gln Gly Asn Leu Xaa Arg Xaa
 1           5           10          15
Cys Ile Xaa Xaa Leu Cys Asn Lys Xaa Xaa Ala Arg Xaa Val Phe Xaa
          20          25          30
Asn Asp Pro Xaa Thr Asp Tyr Phe Tyr Pro Lys Tyr
          35          40

<210> 20
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<400> 20
Ala Gly Ser Tyr Leu Leu Xaa Xaa Leu Phe Xaa Gly Asn Leu Xaa Lys
 1           5           10          15
Xaa Cys Tyr Xaa Xaa Ile Cys Val Tyr Xaa Xaa Ala Arg Xaa Val Phe
          20          25          30
Xaa Asn Xaa Val Val Thr Asp Xaa Phe Trp Arg Arg Tyr
          35          40          45

<210> 21
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<400> 21
Ala Gly Ser Tyr Leu Leu Xaa Xaa Leu Phe Xaa Gly His Leu Xaa Lys
 1           5           10          15
Lys Cys Trp Xaa Xaa Ile Cys Val Tyr Xaa Xaa Ala Arg Xaa Val Phe
          20          25          30
Xaa Asp Asp Xaa Thr Thr Asp Xaa Phe Trp Arg Thr Tyr
          35          40          45

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